RAD21 gene

RAD21 cohesin complex component

Normal Function

The *RAD21* gene provides instructions for making a protein that is involved in regulating the structure and organization of chromosomes during cell division.

Before cells divide, they must copy all of their chromosomes. The copied DNA from each chromosome is arranged into two identical structures, called sister chromatids, which are attached to one another during the early stages of cell division. The RAD21 protein is part of a protein group called the cohesin complex that holds the sister chromatids together.

Researchers believe that the RAD21 protein, as a structural component of the cohesin complex, also plays important roles in stabilizing cells' genetic information, repairing damaged DNA, and regulating the activity of certain genes that are essential for normal development.

Health Conditions Related to Genetic Changes

Cornelia de Lange syndrome

At least six mutations in the *RAD21* gene have been identified in people with Cornelia de Lange syndrome, a developmental disorder that affects many parts of the body. Mutations in this gene appear to be an uncommon cause of this condition.

Some cases of Cornelia de Lange syndrome have resulted from a deletion that removes a segment of DNA on chromosome 21 including the *RAD21* gene. In these cases, the entire gene is missing from one copy of the chromosome in each cell, so cells produce a reduced amount of RAD21 protein. In other cases, the condition is caused by mutations within the gene that impair or eliminate the function of the RAD21 protein. A defective or missing RAD21 protein likely alters the activity of the cohesin complex, impairing its ability to regulate genes that are critical for normal development. Although researchers do not fully understand how these changes cause Cornelia de Lange syndrome, they suspect that altered gene regulation probably underlies many of the developmental problems characteristic of the condition.

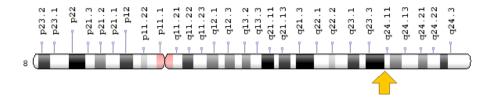
Studies suggest that mutations in the *RAD21* gene cause a form of Cornelia de Lange syndrome with relatively mild features. Compared to mutations in the *NIPBL* gene, which are the most common known cause of the disorder, *RAD21* gene

mutations cause less significant delays in development and growth and are less likely to cause major birth defects.

Chromosomal Location

Cytogenetic Location: 8q24.11, which is the long (q) arm of chromosome 8 at position 24.11

Molecular Location: base pairs 116,845,934 to 116,874,866 on chromosome 8 (Homo sapiens Annotation Release 108, GRCh38.p7) (NCBI)



Credit: Genome Decoration Page/NCBI

Other Names for This Gene

- CDLS4
- double-strand-break repair protein rad21 homolog
- hHR21
- HR21
- HRAD21
- KIAA0078
- kleisin
- MCD1
- nuclear matrix protein 1
- NXP-1
- NXP1
- RAD21 homolog
- SCC1
- SCC1 homolog
- sister chromatid cohesion 1

Additional Information & Resources

Educational Resources

 Molecular Biology of the Cell (fourth edition, 2002): Cohesins and Condensins Help Configure Replicated Chromosomes for Segregation https://www.ncbi.nlm.nih.gov/books/NBK26931/#A3334

GeneReviews

 Cornelia de Lange Syndrome https://www.ncbi.nlm.nih.gov/books/NBK1104

Scientific Articles on PubMed

 PubMed https://www.ncbi.nlm.nih.gov/pubmed?term=%28RAD21%5BTI%5D%29+AND +english%5BIa%5D+AND+human%5Bmh%5D

OMIM

 RAD21, S. POMBE, HOMOLOG OF http://omim.org/entry/606462

Research Resources

- Atlas of Genetics and Cytogenetics in Oncology and Haematology http://atlasgeneticsoncology.org/Genes/GC_RAD21.html
- ClinVar https://www.ncbi.nlm.nih.gov/clinvar?term=RAD21%5Bgene%5D
- HGNC Gene Family: Cohesin complex http://www.genenames.org/cgi-bin/genefamilies/set/1060
- HGNC Gene Symbol Report http://www.genenames.org/cgi-bin/gene_symbol_report?q=data/ hgnc_data.php&hgnc_id=9811
- NCBI Gene https://www.ncbi.nlm.nih.gov/gene/5885
- UniProt http://www.uniprot.org/uniprot/O60216

Sources for This Summary

- Deardorff MA, Wilde JJ, Albrecht M, Dickinson E, Tennstedt S, Braunholz D, Mönnich M, Yan Y, Xu W, Gil-Rodríguez MC, Clark D, Hakonarson H, Halbach S, Michelis LD, Rampuria A, Rossier E, Spranger S, Van Maldergem L, Lynch SA, Gillessen-Kaesbach G, Lüdecke HJ, Ramsay RG, McKay MJ, Krantz ID, Xu H, Horsfield JA, Kaiser FJ. RAD21 mutations cause a human cohesinopathy. Am J Hum Genet. 2012 Jun 8;90(6):1014-27. doi: 10.1016/j.ajhg.2012.04.019. Epub 2012 May 24.
 - Citation on PubMed: https://www.ncbi.nlm.nih.gov/pubmed/22633399
 Free article on PubMed Central: https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3370273/
- McKay MJ, Troelstra C, van der Spek P, Kanaar R, Smit B, Hagemeijer A, Bootsma D, Hoeijmakers JH. Sequence conservation of the rad21 Schizosaccharomyces pombe DNA double-strand break repair gene in human and mouse. Genomics. 1996 Sep 1;36(2):305-15.
 Citation on PubMed: https://www.ncbi.nlm.nih.gov/pubmed/8812457
- Minor A, Shinawi M, Hogue JS, Vineyard M, Hamlin DR, Tan C, Donato K, Wysinger L, Botes S, Das S, Del Gaudio D. Two novel RAD21 mutations in patients with mild Cornelia de Lange syndrome-like presentation and report of the first familial case. Gene. 2014 Mar 10;537(2):279-84. doi: 10.1016/j.gene.2013.12.045. Epub 2013 Dec 27.
 Citation on PubMed: https://www.ncbi.nlm.nih.gov/pubmed/24378232
- OMIM: RAD21, S. POMBE, HOMOLOG OF http://omim.org/entry/606462
- Sonoda E, Matsusaka T, Morrison C, Vagnarelli P, Hoshi O, Ushiki T, Nojima K, Fukagawa T, Waizenegger IC, Peters JM, Earnshaw WC, Takeda S. Scc1/Rad21/Mcd1 is required for sister chromatid cohesion and kinetochore function in vertebrate cells. Dev Cell. 2001 Dec;1(6):759-70. Citation on PubMed: https://www.ncbi.nlm.nih.gov/pubmed/11740938

Reprinted from Genetics Home Reference:

https://ghr.nlm.nih.gov/gene/RAD21

Reviewed: September 2015 Published: March 21, 2017

Lister Hill National Center for Biomedical Communications U.S. National Library of Medicine National Institutes of Health Department of Health & Human Services